3rd Bay Area Population Genomics Conference

Sponsored by the Ecology and Evolution Group at the Biology Department at Stanford University

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Registration is FREE but REQUIRED

Date:	December 11, 2010 (Saturday)		
Place:	tanford University ~ Biology Department ~ Herrin Hall ~ T175 Auditorium ~ 393 Serra Mall ~ Stanford, CA		
Sign up RSVP by Dec. 6	Attendee List (current count: 99)		
Directions	Searchable map, 393 Serra Mall, Herrin Hall		
& Parking	FREE Parking since it's a weekend \sim Available at Lomita Drive, Palm Drive, Museum Way, the Oval		
Contact:	Dmitri Petrov, Stanford; <u>dpetrov@stanford.edu</u> Joanna Kelley, Stanford; j <u>okelley@stanford.edu</u>		

SCHEDULE

	9:00 AM	Breakfast Buffet	
	9:30 AM	Rachel Brem, UC Berkeley	Pathway evolution in Saccharomyces
	10:00 AM	Dario Valenzano, Stanford	Genetic Architecture of longevity in the short-lived fish Nothobranchius furzeri
	10:30 AM	Paul Jenkins, UC Berkeley	A new approach to computing likelihoods in population genetics models with recombination
	11:00 AM	Coffee Break	
	11:30 AM	Jared Wenger, Stanford	Adaptive mutations effect minimal trade-offs across the yeast adaptive landscape
	12:00 PM	Ed Green, UC Santa Cruz	Recent human evolution as revealed by ancient hominin genome sequences
12:30 PM Lunch and poster session (register if you would like to get food!)			ou would like to get food!)
		POSTERS (signup is now closed):	
		Barbara Dunn, Gavin Sherlock Lab, Stanford	Analysis of the Saccharomyces Pan-Genome Across Diverse Strains Reveals Genome Plasticity in Response to Industrial Environments
		Diamantis Sellis, Petrov Lab, Stanford	Fitness overdominance as a generic feature of adaptation in diploids
		Vera Kaiser, Doris Bachtrog Lab, UC Berkeley	Non-random gene loss from the Drosophila miranda neoY chromosome
		Dan Kvitek, Gavin Sherlock Lab, Stanford	Sign epistasis between adaptive mutations causes a rugged fitness landscape during experimental evolution
		Beatriz Vicoso, Bachtrog Lab, UC Berkeley	Lack of global dosage compensation in Schistosoma mansoni, a female- heterogametic parasite, Doris Bachtrog Lab, Berkeley
		Alan Bergland, Petrov Lab, Stanford	Natural mutations in the aquaporin, Drip, affect Drosophila fecundity
		Fabian Staubach, Petrov Lab, Stanford	Is there ancient admixture between house mouse subspecies?
		JJ Emerson, Doris Bachtrog Lab, Berkeley	Fisher's geometrical model in light of experimental evolution data
		David Goode, Arend Sidow Lab, Stanford	Assessing the functional impact and selective constraints acting on short indels in the human genome

Kirk Lohmueller, Nielsen Lab, Berkeley	Natural selection affects multiple aspects of genetic variation at putatively neutral sites across the human genome
Matthias Steinruecken & Josh Paul, Yun Song Lab, Berkeley	An accurate sequentially Markov conditional sampling distribution
Michael Palmer, Feldman Lab, Stanford	The Red Queen's Race can select for evolvability
Depart or go for a hike around Stanford	

2:00 PM Depart or go for a hike around Stanford

<u>Eco / Evo</u>

<u>BAPG Google</u> <u>Group</u>

<u> Map</u>

<u>Biology</u> Department **Stanford**